

# POLYMORPHISMS IN THE TNFRSF1A GENE

CGGACATAGC	CAGATGTATT	ACGGATGACT	GCAGTCAGCT	CCCCCAGGCT	
CCTGCTTCTC	TTGCCTCCTG	CTTTTTTCCC	CAGAGCTGTC	TCCTTATCTC	100
CATTCACTTG	TCTATGGGTT	ACTCCTGGAC	CCTGGGGTTA	GGAGTTGGAA	
TCAGGCTGTT	AGCGATAAAA	GGGTTC AAGT	TGACTCATT	TCCTTATCAG	200
GCTTAGTAGT	TGAAGTGACT	TGCTGAGCTT	CATAATTCTT	AGAGAACCTG	
CCATGAACCC	AGCTCCCTTT	CTATGACTCA	CCCTGCCACC	CTGTGACACA	300
TAGAGTCTGA	ATGGCAGGTC	TGGGGCTAGA	ACCCACGTCA	TCTGGACTTG	
GAGTCCAGTG	ACCCTTTGGG	TTAAGCATGT	GTGTGTGTGT	GTGTGTGCCA	400
TGATGCGGGA	GGAAGGTCCC	TGCTCTCTGT	AGCTGTTTTT	TTTCATCCTTT	
GCTCTACAAG	CCCTAACAGC	CGATTCTGTC	ATCCCTAGTC	TGCCCCCTCTC	500
CTGTTTCTCC	ATCTCCTCTG	ACCATGATTT	TTTTCTGTCC	CTGGAGGGAT	
GATGGTCTCA	TTCTCACCTC	CTCCACGAAA	CGTGTTAGCT	TTTCATATTC	600
CTAGATCCAC	TCATTCTCA	TCATCTTTTT	TTTTAAACAA	AATTTTATTG	
AAAAATGTAA	TATGACGTGT	CAAAGTTGTA	AAGTTATTGA	GTAAATAAGC	700
ATGTATCCTA	AATATTGAAA	AATATTCTCC	TTTTGTACCA	GGCTATGTGT	
CACGGCTTTG	GCGCTTTGCA	CAGACTATTA	GAAATACCTT	ATAACATTAA	800
AAATAGGACA	TTGAGGCCGG	GCGTGGTGGC	TCATGCCTGT	AATCCCAGCA	
CTTTGGGAGG	CCAGGGTGGG	TGGATCACCT	GAAGTCAGGA	GTTTGAGACC	900
AGCCTGGCTA	ACACGGTGAA	ACCCCGTCTC	TACTAAATAC	AAAAAATTAG	
CCGGGCATGA	TGGCACATGC	CTATAATCCT	AGCTACTCGG	GAGGCTGAGG	1000
CAGGAGAATT	GCTTGAATCC	GGGAGTCAGA	GGTTGCAGTG	AGCCGAGATT	
GTGCCACTGC	ACTTCAGCCT	GGGCAACAAG	AGTGAAACTC	TATCAAAAAA	1100
AAAAATAGGA	CATTGAAGTT	GGTTTCTTTT	TTTGATACAG	AGTCTCGCTC	
TGTCACCCAG	GCTGGAGTGC	ACTGGCAGGA	TCTCGGCTCA	CTGCAACCTC	1200
TGCCCTCCTGG	GTTCAAGCAA	TTCTCCTGCC	TCAGCCTCCT	GAGTAGCTGG	
GATTACAGGC	ACGCGCCACC	ACGCCTGGCT	AATTTTGTAT	ATTTAGTAGA	1300
GACAGGGTTT	CACCATGTTG	GTCAGGTTGG	TCTCGAACTC	CTGACCTTGT	
GATCCGCCCA	CCTCAGCCTC	CCAAAGTGCT	GGGATTGCAG	GCGTGAGCCA	1400
CCGCACTCTG	CTTTTTTTTT	TTTTTTTTTG	CGCCCTCTCA	CATACCATAC	
TCCCCTGTAT	CACCTATCCT	TCTGAAGTTG	TTATTAATCA	TTAATACAAC	1500
TAGCTGGGCA	TAGTGGTGTG	CGATGGTAGT	CTTAGCCACT	CGGAAGGCTG	
ATGTGGGAGG	CTAGCTTGAG	GCCAGTAGTT	CTAGGTTAGG	TGAGCTATGA	1600
TTGCACCATT	GCACTTTAGC	CTGGGTGAGA	GCAAGCTCCT	GTTTCAAAAA	
AAAAATTAAT	TGCTACCACT	TACTAAATGC	TTAATATATG	GCAAACACTT	1700
GCCAAACACT	TTATATGCTT	GATTTAAGCA	TCAAGCTAGC	TCTGTGAAGG	
GTACCAGCAG	GTTTCCCATT	TTTTAGATGA	GCAGACCGAG	GTTCTTCTCG	1800
CTGCTTCATA	CTGGAAACTT	GCACTTGATT	CTGAGGCTCC	TGCTTCTTCA	
AGAACACTGC	TTTGGGTTCC	CTTCTCCTGT	CCCTGGGGTC	TCCCTTTGTG	1900
ATGGTGGTGA	GCTGCTTCCT	TTCTGAATCC	AGCTTCAACC	CTACAGTTCT	
CCAGAAGCTG	GACGATGGGG	TGGAGTAAAG	TCAGCTCCCC	CCGCAGTGAG	2000
GGACACTGAA	GCTCCATTCT	CATCTGCGGA	TCACAGAGGG	GAAGCCAGGA	
AGAGCCAGGG	GACGGTGGAC	TTGGGGCTGG	GAGGTCATCT	CAGAGGGATA	2100
AGGGGTGAGG	AGCTCTGGTT	TCAAGTTCCA	AAGCCCTAGG	ACCTCCCTCT	
TCTCTGTCTG	CCTGCATTTT	TAGCAGCCTC	AGCAGCTGCA	GGCCCTTGGG	2200
CGGGGCTGGA	TGTAGGGAAG	GTCATTGTAC	CAAGAAGATA	GTTGGGTAAA	
TGTGGTACCT	TTGTTGTAGG	ATTCTCTTGG	GAGATGTCTG	CATCAATGAG	2300
GATGGCATAA	AGTAACCAGA	GTCAGGATGT	GGGGTCTGAC	TCAGTGACAG	
AAAAAGTGGC	AGTGTGTCTC	TCATAGCCAA	AGGGGCCCTT	GGACCGGCAG	2400
TCGGGAGTCT	GGGGTTCTCT	GTTGGCTCTG	CCTCCTGGCA	CATTGGGTTT	
CTGGACCTCA	GTTTTCTCCT	CTATAAAACC	GGGCAGTTGG	GTGGGCACGG	2500
TGGCTCACAC	CTGTAATCCT	AGCACTTTAG	GAGGCTGAGG	TGGGCAGATC	
ATTTGGGCCC	AGGAGTTCAA	GACCTGCCTG	TGTAACATGG	TGAGACCCTG	2600
TCTCTACAAA	AAATACAAAA	ATTACCCAGG	CGTGGTGGA	TGCACCTATA	
GTCCAGCTG	CTTGGGAGGC	TGAGGTGGGA	GGATTACTTG	AACCTGGGAG	2700

FIGURE 1A

GTCGAGGCTG	CAGTGAGCTG	CGATGGTACC	ACTGCACTCC	AGCCTGGGAA	
ACGGAGCGGA	CCCTCAAAAC	AAAAACAAAA	ATGAAAAACA	AGCAAACGAA	2800
GAAATAAAAA	AACCTAGGGG	GTTGTAGTCG	ATGATCTGTA	AGGTGAGTTA	
TAATTGATGT	ATTGGAATAT	TTAGGAAAAG	GGCACTGGGA	ATATGCTAGG	2900
AACACCTGAT	GGAGGTATCT	TTATTTCCAC	GGCAGCTTCG	TGGATACGTC	
TCATTGATTG	TCATGGCATC	ACTTTCCCCA	TGTAGGTGGG	CAGACATTGT	3000
TACCCCTGTT	TAATAAACAA	GGAACCAACA	GAGGCTTAGG	AGAGGAGTTG	
CCTGATGTCG	CATGATTGGT	GGCAGAGCCA	GGATCAACAG	TGGGGCAGGG	3100
TGGGGGGACC	TGGCCAGGCA	GAGACTGGAT	GAGACCTGGG	GTGAGGAATG	
T					
GCAGGCACCC	AGTCAGGGCA	GAAAACGAGG	GTTGGGACTT	ACTTTGAGTT	3200
TTGGATTGGA	TCAGTAAATT	CCCAAGAAAG	AGGGAGACTA	GGAGGCTAGT	
GAAGAACTCT	GGAGTAAAGG	GGAGGATTAC	TAAGGGACAT	GGAGTACCTA	3300
TCATGTGTCTG	GACGCTTATC	TATATCTCTC	CCATCTGAAC	AAATCCTTAC	
AGGAACCCCA	GGAGACAGGT	TATCTCCACT	CTGCAAATTG	GAAAACAGAT	3400
CCAGACAGTT	TCAGTTATGT	GTCTGAGAAG	TTCATTTATG	TGTCCAAGAC	
G					
ACATTCTTAG	CTAAAAAGCT	AAGCATTCTG	AATTGGAACC	CAGAGAATTT	3500
GACTCCCAGA	CTCTGGATCT	TTTCACTGCT	GTGATCCATC	TGGGAAAGGC	
TAGTGATGTG	GGCAAGGGGC	TTATTGCCCC	TTGGTGTTTG	GTTGGGAGTG	3600
GTCGGATTGG	TGGGTTGGGG	GCACAAGGCA	GCCAGATCTG	GGAATCCTGT	
G					
GCTTGTGACT	GGACTACAAA	GAGTTAAAGA	ACGTTGGGCC	TCCTCCTCCC	3700
GCCTCCTGTG	GCCTCCTCCT	CCAGCTCTTC	CTGTCCCGCT	GTTGCAACAC	
TGCTCCTACTC	TTCCCCTCCC	ACCTTCTCTC	CCCTCCTCTC	TGCTTTAATT	3800
TTCTCAGAAT	TCTCTGGACT	GAGGCTCCAG	TTCTGGCCTT	TGGGGTTCAA	
GATCACTGGG	ACCAGGCCGT	GATCTCTATG	CCCAGTCTC	AACCCTCAAC	3900
TGTCACCCCA	AGGCACCTGG	GACGTCTCTG	ACAGACCGAG	TCCCGGGAAG	
CCCCAGCACT	GCCGCTGCCA	CACTGCCCTG	AGCCCAAATG	GGGGAGTGAG	4000
AGGCCATAGC	TGTCTGGCAT	GGGCCTCTCC	ACCGTGCCTG	ACCTGCTGCT	
[EXON 1: 4019..					
GCCACTGGTG	AGACCAGGGA	CAAAGGGAAG	AGTGGGCTGG	TGGGCGAGGC	4100
G					
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ACCTTCCGGC	TGGCGTGGGC	CCTCTCCGGG	AGGGGGCCGA	GCCTCTCCTG	
CCCGGGCCTG	GTCCTGGCGC	CAGCCTCAGG	CCTGCAGGTC	CTAACCTCAG	4200
CCACTGCCAG	TGTGGGGTTC	CCCATTTCATC	CGCCTTTTGG	AGTAGGGGCT	
GCGCTGAGGC	AGGGGAATGG	GAGAAGTTTG	AAAGGGAGAG	AGTAAAAGGA	4300
AGCCCTGGCC	CCTGACAGCG	GTGGAAGTTT	GTGGGCGGCC	AAGGGAATGT	
GGGCAGGAGA	TAGGCCCAGG	GTGGGGCAGA	TTTGGCGGGG	AAAAGAAGGG	4400
AGTGGGAGTA	GGAAGATTAG	TGCTCGGGGA	GTCCAGACGG	TTCTGAATTC	
TGTCCCTCCG	GTCAGCTGGC	TGGCCTGGAG	GGTGTGGGGC	CGTGGGGAGG	4500
CGAGGCTGCC	TGTGGAACCT	GGTGGAGCAC	ACCCTGTAGG	GCAGGATTTT	
GGCGGCTGGT	GAAGTGGGGG	AGTGAGTTGA	GGAGTGGGGA	TGGGCTGGTG	4600
TGGTGGGTTT	GGGATGCTCA	TGGTGGGAGG	TATTTGAGAA	TGGGCTGGGA	
CACTGGATGG	GGCAGGGCAA	CCCAGTGGAC	AGTGTCCCCA	GTGCCCTGGC	4700
CAAGCCCCGG	CCTCTCACCT	GGGGACATTC	TTTACCCTTT	TGCCTGCTGC	
TAGGCAGGTA	GCCGCTGTGG	GA CTGAGCCT	TCCCAGGGAG	CTAGTCCTAC	4800
CCCCACCTGG	TCAGTGTCCC	TGGGCCTGTC	CTCCAGCTTC	CCCTCCCCGC	
TGCTTCTCAC	AGACCTAAAC	AACAATCCCT	TGGTTTCTTA	TTCTACAGTT	4900
CAGTTTGGGG	AAGTTGGTAG	AAAGTTGTTT	TCGTCACTGG	AAAATGTCCC	
TTTCTCTGGC	CTCAGCCTTG	TTTCAATGTA	TCCTTGATCG	TCCTCCACGT	5000
CTTGGTCCGG	GAATCATCCT	GTTTCAATGT	CCTGGGCCCA	TCTAGTCAGG	
CAGATTTTCC	CTGCCCTGCC	CGGCCTCTGA	AGGCTGCGCC	TACCTCCCCT	5100
CTCTTTAGTG	CCTTATACTC	TTCTCTCTCT	ACCATTCTTT	TCTTCCAGCA	
ATCTCCCCAG	ACTCTCCTCA	GACTTCTCTAG	AGCCTCTTTT	TTTGAAATCT	5200

FIGURE 1B

TTTCTCGCTA	ATCCTCCTTC	CCCTCCTCTC	TGCTCCGCTC	TGGTCCCGGC	
CCCAGGTCCC	CAGGCAGCAC	GTCTCTGGTC	AGGGTCTCAC	TCTTCTTCTT	5300
CTGCCTCCTC	CTGCCTCCTT	AGTCCCACCC	GCTCTTCCCT	TCTTCCCCT	
GTCTTTCCCC	CACGGTCTCC	CCACCAGCCA	GCTGCCCTGA	CATCCTGCTT	5400
CTGTTTTCTG	TTTGGGGGCG	GCCCCTGGCT	CCCTCACATA	CCTCCTGCAT	
GAACAAGAGC	AGCTTATATA	ACCTAACCTT	CCATGCCTTC	GTTTCTTTAT	5500
CTCCAAAATG	GGTGTACACAG	TCTTGACCTC	ATACTGTTGT	TTTGAAGATT	
GAATAGACTG	ATACATGTTA	AGTGTTCATT	TGATTTATTA	AGTGTGCGCT	5600
CTGGGCTAGA	CACTGTGATA	GGTGTGGGA	TTACAGCAGA	GAACAAAATC	
CCTGCCCACA	GCTTTGACAG	TCCATCAGGG	GAATAGGTTG	TAGCAAATAG	5700
AAAGCACTCA	ATAAAGTTTT	TATATTGCTG	TGACTAGTAG	TAATTACTGG	
GTGGCTACCT	GTGTTGGGAA	AACAGAGGGT	AAAGGTAGCC	TGAACAGGTA	5800
AAGGGAAGTG	CCTGCGTCCT	GGGGTGCTTC	AGCCCAGGTG	GGATTATGTC	
TCCTAAGGGA	CAGAAGCCTG	GCCTGGAGCT	GGAGGAAAGG	GAAAACAAAG	5900
GGAATGCAAC	ATCCTTCTGA	ATTTCTCACC	ATTCAGTGGG	CAATGCAGAG	
CTCACAGTGT	GTGTGTGTGT	GTGTGTGTGT	GTGTGTGTGT	GAGAGAGAGA	6000
GAGAGAGAGA	GAGAGAAGTG	GGGTAGGGGA	GTAGGGGAAGA	ATGATACAGG	
AGAGACTGTG	GCAAAGCAAA	CAGGATTTTG	CTGCTCTCAA	AGAGCTTACA	6100
GCCTAGTAAC	CAAGATGGCT	TACAGTGAAA	AATGATTTC	GAGCAATCCC	
GAGGAAAATA	TCCACAAATG	CATTGTGATG	TGGTGTCTCT	GAGCACCAGT	6200
TGGGAGGAGG	AGGAACTGGG	GAAGGAGGTG	AGCCTTAGTC	CACTGCCTTT	
CCTTGCTTAG	CAGGTCTCAG	CTCCTGCGCT	CAGCTCCAGA	AAATTCAGGA	6300
GCTTCCCCAC	GCTGCTTCAG	TGTCCTTCAC	TGTGCAACTG	CAGCACTCCC	
TGTATAGATC	TCAGTGCCTA	CAACTGACTG	TCTTTGACTC	AAGTGAGAGC	6400
TCTTGAGAGC	ACGAGCTGTG	TATTATCCAC	CTCAGCATCC	CTAGCACCCA	
TACGGGACCT	GTACACATTAA	CTGTGCCCTT	TAACATTTTG	CTGAAGGAAT	6500
TAAGGAACAA	GAGATGTGTC	AGATGGGATG	GCGGAGGGAA	AGCCTCATAG	
AAAAGTGGAT	GTGGAGCTGA	CATCTGAAGT	CACTGCCTGT	CAGGGTAGCT	6600
ATAAAGGAGG	GAAGCAGAGT	TGGATACTGA	TGTGAGGAAG	AGGAGAGGAA	
TGGAGAGATG	GGATTTTGTG	TTGATGGGCA	GGGTGGCAGG	AAGCCAGACA	6700
CCTTGTTTCG	GGAGTGGAAA	AACCATGTTG	AGAAACACTA	AGAAATGTGA	
ATGGGAGAAT	TAGAGGGAGT	GGGGGAGAGG	ATGGAGGAAG	AGTGTGAAT	6800
ATGGTTCCAG	GTGGAGGAAT	TCATTCAATC	GTTTATTCAG	AAGCTGTTCT	
CCTAGGGCAC	ATTCTGTGCC	CAGACTGTGA	TTAGAAGTGA	GGTGAGGCAT	6900
CTCAGATGGG	TGCTGTGGTT	CATGCCTGTA	ATTCCAGCAC	TTCAGGAGGC	
CGAGGTGTGT	GGATTGCTTG	AGTCCAGGAG	TTGAGAGCCA	GCCTGGGCAA	7000
CACAGCAAAA	CCCTGTCTCT	ACAAAAAATA	CAAAGATTAG	CGGGGCATGG	
TGGGGCGTGC	TTGTCATCCC	AGCTATTCGG	GAGACTGAGC	TCGGGAGGAC	7100
GGCTTGGGCC	CAGGAGGTGG	AGGTTGTAGT	GAGCCCTGAC	CACACCACTA	
CATTCCGTCC	TGGTGGTGAA	GGTTGCAGTG	AGCTATGATT	GTGCCACTGC	7200
ACTTCACCTT	GGGTGACAGA	GTGAGACCTT	GTTTCAAAAA	AAAAAAAAAA	
AAAGTAGTGA	GGCATCTGTG	GAAGTCTTCA	GATCATTTCC	ATGACCATGG	7300
AAATGCTGTT	TGGAGCCAGG	CCCTGGAGAT	GGAGAGGAAG	GTTACACAC	
TTGTGCGTGC	AAGTTAAAGC	CTGAATGAAG	ATTTAAAAAG	TGTGTAGGAC	7400
GGATGGGAGC	AGGAGAGAGG	CTAGAAGACA	CTTGCAATAA	CCCAGGTGTG	
AGGCAACCCA	GGAATGCGGA	GAGGACCGAG	AGATCACAGG	GGGAGGCCTC	7500
GCAAGATGAA	CTGACACATG	GGATGGCGGC	AGGGATAGGG	ATGGGGCCCT	
GGGGAGAGAG	CGTGGCAAGT	TCTCAGCATT	CGTCCGGGAA	GTCGATGGTG	7600
TGTCATTTGT	CTAGGTGAGG	AGATGGATGA	ATTCCGTCTG	GGGCATGTTA	
AGGGTCAGGG	AAATGGTCAT	GTGGAAGGGT	GCGCCTACCA	AGCTGGAGGA	7700
GAGGTGCTGC	AACTTCTTTC	TGCCTTTGTA	TCATTACAGC	ACACTGTGTT	
CACTCATCAG	TGGTTCTCAA	AAGGAGAGGA	GCACACCAGA	CTCTTAAGTA	7800
AGGGTGTGTG	TGCTTGTGTG	TGGGGAGGTG	GGGGGATGGT	CTGAAAACCTC	
TCCCCCGGAG	ATAAATATAT	TCCTACCAGG	GGTGCTGTCT	CCTCACCTCC	7900
CTCTTTGGGA	ATCACTGGCT	TCTACTAGAG	TGGAAGACAG	ATGTATCATT	
AGATCGATCA	GTTGATCCAT	ATTTATCTGC	TCCAGTCTG	GAGGTCTGGT	8000

FIGURE 1C

TCTGGGAGCT	GAGAGGACAC	CAGGGGAGGA	TAAGACACTT	TCTGACCAAG	
ACATTTTTTTG	ATCTCTCATC	TTATAAGGTT	CGTGGTCACT	TTGGGGAGAT	8100
CATATCTGTC	ACCCAACATA	ACCATATTAT	GATAAGAGCC	AAAAGTAGAT	
AGGGTCAGTT	CACGTGCTTC	GAGTTCACAG	GGACTATGGG	TCTAAGGAGC	8200
CGGGGTGGAG	GAAACAGACA	TCGTCAATGG	TGGCTTCACG	GGAGGGAGAT	
GGGATCTCAA	CTGGGCCCTT	GGAGGAGAAG	CTGCCACGAC	CTCCCCCAAC	8300
ACCTTGACAT	TAAATGAACA	GACACATGAA	TGAGGGGGAA	AGGAAGACTA	
ATTGGGTCCC	TGCAAGGTGG	CTGGATCGGG	GTCAGACCAC	AAGGCCGATC	8400
TCAGCGTCGC	CTCCCCACTC	TGCAGCCCCA	GCACAGGAAG	TCACACTTTA	
AAGCCTCCTC	TGGCGGAAAT	TGTGGGGGAG	TTGGAGGGGT	GTTGGGCCAC	8500
CCCCTCAACT	GTCTCTCCAC	AGGCACCCCA	GCTTCCTGCC	CTTCTGCTCC	
AGGCTGGAGT	CTGGGCCTAA	AGAGCTCACC	TCCTGTTTCT	CCTGTTTTGC	8600
TTCATTTACG	CAACTGCTGA	GGACTGGGCT	TACTGGGGCC	AGCTGGTGCC	
AGCAGTGGTG	CCCAGTGGTG	GGGAGTCTGA	GGGCCCTGGC	TCCTAGGGAT	8700
CAGAGAGGGC	TGACCTGGAG	CATTCTGGGG	GCCAGGGGAA	GCCTAGGAAG	
CAGGGCTGGT	TCTTCCATCC	GGCATCCCTT	CTTGCCTGCT	CCCTCGTTCC	8800
TGGAAGTGGG	TGTTCAAGGC	TCTGGAGGCT	TTCTGTATT	GCCAGTGGGC	
TTGGGGAGGG	TCTGTGGAGA	CTCAGAACTG	GCCTTGTTTC	CTAAGGATTG	8900
TCTGGGGACC	CCAGGGAGGC	CCCCAAACCC	AGCACAACTG	GTCAGAACCA	
GCCAGGCTGT	GGGAATGCGG	TGAACCCAGG	GTGGGAGGGC	AGCCTTGCT	9000
TGCTTCCTGC	TGGGACTGGG	GAGTGTGGG	GGATGGAGTG	AGAGCTCACG	
GAATGGGTTT	AGCTGTTGGA	GACTTGTTGA	ACTGGGAGGA	GGAGCTGGGG	9100
CGGGGCCCTCA	GCTAAAGGCC	GCTGAGGGGC	TAGGAGGAGC	CAAGTGGCCC	
TCAGGGGAAGG	GAGGGCACAG	ACCTGATGGG	CGGAAGCCAG	GGTCGAGGGA	9200
GACTTCCCTT	CGGGATGGAA	TGGGGAGAGG	GAGGCATTTC	CCGGAACATG	
TGGGCCAAGT	GGGACAAGGG	TCTGTGGCCT	GGCTCTTTGC	ATGGGGAGGG	9300
GATGGATGGG	GGTTGAGTGG	GGATGGGAAG	GAGGGACTTG	GCCATAGGAA	
GAAGGGATTA	GATGGAGTCC	CACCTGTCATG	CAGGCTGGTG	CCTTCTGCCT	9400
TTCTGCTGAC	TCATGACCCT	TGAGGAGCTG	GGGAAGCTGC	TAGTTCCCTC	
TCCCCCTCCCT	AGGTCTCCCT	CCCTCTGGCC	TGAGTCACTG	GGGCGGAGTT	9500
GCTGGGAAAA	GATTTCCCTT	TCCCGGATCT	GACTTAACCC	CCAGAGTGCT	
GGAAAGAGAA	GGGAACACGT	GGCCTGAGAA	AGCCTCTCTC	CCTCCCTCCC	9600
TCCAGGGAGG	CTCATCCCCC	ACTGGCCAGA	GGTCCCTGAA	AAGCTCCCTT	
TAAGGCTGTC	TGGGGCTGGC	GTCCCCCAGT	TCTTCATCAT	GACTCTGCCT	9700
CAAGCCCCCT	GGATGGGATT	CAAAGTACCA	GTGACCTTAG	GTGCTCCAGT	
GGCTTCTTCG	GGGAAAGGAA	CCACACTTTC	AGGACTGGGA	AGTTCTTCCC	9800
ATCACCACCC	CAAACCCTTC	CTGTTGCCCT	GGAAGCCCCA	GTCTGTCTCT	
CAGCAGAGGT	GGCACGGTGT	TGGCTGGTGC	GGGCAGGGGA	AGGTTGTTGT	9900
CCTCTGAGCA	GGGGCACACG	CCTCCACCTG	CGGGGGCTGC	TGTTGTGTTT	
CTGTGTGTGG	CTTCCCCTGT	TTGCGGCTGA	GGCTTGAAC	TCCGGGCCTG	10000
CACAGCTTAC	AGCTGCAGCG	TCTCCCCGTG	GCTGACTCAG	GGTGACTGGC	
CTCCTGCTCC	GAAATGTGGA	GTTGGTGAGG	CTGGGTGGCT	GTGGGCTGCC	10100
TGACCCTCCT	TCCCTGCCCT	AGGGTTTCTG	TGATCTGGTG	AGTCAGTTGC	
TCCCCAGTGT	TTAACAGACA	TTGAGGACAC	CCTCTTATCT	TTACACAAAG	10200
TGTCTCTTAT	AGTAGAAAAA	AAAAATGAAG	CCCAGGGAAA	ACCAGAAATG	
AAGCTGGCAG	AGATCAAAGT	CCAAGTTAGA	GCTAAATATT	CACTCCTGGC	10300
TTTGCTTTCC	TGGCACTGAT	GCCGGAACAG	GACAAGCCAT	TTAGCTGCTG	
TGGGGTTGGC	CTGAGACTGC	AAAGCACACC	TTCCAGAATG	CCATGGTGTG	10400
CAGGGGGCTC	CAGGACTCCC	CAGCACGCCC	TCAGCTCTGA	CCTGACAGTC	
ATCCAAGCTG	GGTCGCTAGC	CTTGGCCAGC	TCTATTTGCC	TATGTCCTGC	10500
ACACCTTTGC	CCACTCCTGC	CCCCGTCTCA	ACTTTGTCCC	CCGTCTACCC	
ATGCAGGATC	CCCAACCTTT	CCCTTTTACT	CTCCTCCCCA	TTTGTCTTGG	10600
CCAACCCCGG	GTGTTTGTAA	ATTTTGAGGT	GGAGGGGATG	GGCCAGGGAA	
TGTGAGGGCG	GAGGCAGATT	GAGGTTTGAT	ACAACATGT	AAATAAACTT	10700
CCTTCTTCTG	TCCAATCCCC	AGGAGTGGTG	CTCACGGGAA	CATCACTCGC	
CCCCACCGCC	AGCTGACTTT	TTCAGAAAGC	TTTTTCATGGT	GTAACATATT	10800

FIGURE 1D

CCTGGGATGT	GCATAGATCC	TCATTGTTTA	CCTCTGTGAA	TGTTTCGCAA	
GCGATCACAC	GGTGAACCCA	GCACCCAGAT	GGAGAAACAC	CGCCCCAATC	10900
TTTAGGGCTG	CTTGTTGGAA	GAAGGGGCCA	TCCTGAAGT	AACCTGCCAA	
TTCCCAATCA	AAAACACATC	CTTTCAACAT	CTGCCCTGTG	TCCAGCACTG	11000
TTAGCTGCTG	TGGGGGATTT	CACAGTAAGG	ATAAAATACA	GGGCTGGGCT	
CACGCCTGTA	ATCCTAGCAC	TTTGGGAAGC	CAAGGTGGGA	GGATCACTTG	11100
AGCCCAGGAT	TTTGAGACCA	GTCTGAGCAA	CGTAACAAGA	CCCTGCCTCT	
ACTAAAAATA	AAAAAAAATT	AGCTGGGCAT	GGTGGTTCAC	GGCCGTAGTC	11200
CCAGCTATTC	AGGAGGCTAA	GGTGGGAGGA	CTGCTTGAGC	GTGGGTGGTG	
GAGGGTGCAG	TGATTGCATC	ACTGCACTCC	AGCCTGGACA	ACAGAGCAAG	11300
ATCCTGCCTA	AAAAAAAAAA	AATACAGCTT	AGATCTGGGG	CCTACTAGCT	
TTGAGTTGAG	GGAACAAAAA	TGAACACACA	GGACAACTAG	AGAACAATTA	11400
AGCATCAGAT	TGTATGGCCC	CAACTGTCTA	AGTTTCAAGG	AAGAACTCTA	
AACTTAGTGA	GTGGCGTGGC	CTGGGCGGAA	TGTTTCACTG	AGGAAGGACT	11500
TGAGCCAGGG	AAGTTTTAGA	TCTGCTACCC	CTAAGCTTCC	CATCCCTCCC	
TCTCTTGATG	GTGTCTCCTC	TATCTGATTC	TTCCCCAGGT	GCTCCTGGAG	11600
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CTGTTGGTGG	GAATATACCC	CTCAGGGGTT	ATTGGACTGG	TCCCTCACCT	
AGGGGACAGG	GAGAAGAGAG	ATAGTGTGTG	TCCCCAAGGA	AAATATATCC	11700
ACCCTCAAAA	TAATTCGATT	TGCTGTACCA	AGTGCCACAA	AGGTAGGGGC	
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AAGTGGAAC	GGTGAATGCC	CTCAGGTCTG	GGGTGCTGCT	TCTTTCTCTG	11800
CTTCTTCCAG	TTGTTCTTCC	CTAAGTTTGC	TGTCTCTCCT	GGGCTGGGAT	
TTTCTCCCTC	CCTCCTCTCC	TAGAGACTTC	AGGGAATCGG	CCCTGGCTGT	11900
TGTCCTTAGC	ATGGGGCTCC	TTCCTTGTGT	TCTCACCCGC	AGCCTAACTC	
TGCGGCCCCA	TTCACAGGAA	CCTACTTGTA	CAATGACTGT	CCAGGCCCGG	12000
T					
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GGCAGGATAC	GGACTGCAGG	GAGTGTGAGA	GCGGCTCCTT	CACCGCTTCA	
GAAAACCACC	TCAGACACTG	CCTCAGCTGC	TCCAAATGCC	GAAAGGGTGA	12100
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GTGTGCACAG	GCAGGAGAGT	CAGGCGGGTC	TTGAGTGGTG	TGTGGGTGCC	
TGTCTATGTG	CAGGCTGGTG	GGTGTGGGCA	GGAAGGTGTG	TGTTTTGGTG	12200
GGACACTGCA	TGGATGTGAG	TGTGTATTAC	AGAGACACAC	ACTTAGGGGT	
ATGTCAGGAA	GGGGATGCAG	GGACAGGAGG	ATGCAGGACT	CATACCCCAT	12300
CTTCTCCCTT	CACCAGAAAT	GGGTCAGGTG	GAGATCTCTT	CTTGACACAGT	
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GGACCGGGAC	ACCGTGTGTG	GCTGCAGGAA	GAACCAGTAC	CGGCATTATT	12400
A C					
GGAGTGAAAA	CCTTTTCCAG	TGCTTCAATT	GCAGCCTCTG	CCTCAATGGG	
ACCGTGCACC	TCTCCTGTGA	GCGCAGCTCT	CCTGAGGCCA	AGCCCTCTCC	12500
T					
..12466]					
CCACCCCAGG	GGTTGGCCCC	TTCCCCATGC	GGTGGCACTT	CCTTTCTTTC	
CCCCTCCTGT	ATTCTGTGGG	TCTGACAACC	AACTCCTCTC	TGGCCGCCCC	12600
CACCCTGTCC	CTCGTCACTT	CCTCTGTCTT	GTGGGGTGGG	GGTGCAGGCG	
CTTCTCCTTT	AGCTGTGCCG	CACTTCTCCC	TACAGGCCAG	GAGAAACAGA	12700
C					
[EXON 5: 12686..					
ACACCGTGTG	CACCTGCCAT	GCAGGTTTCT	TTCTAAGAGA	AAACGAGTGT	
GTCTCCTGTA	GTAAGTGAGT	ATCTCTGAGA	GCTGCTGGGC	ACTGGATGGT	12800
..12764]					
GGCATGGGTT	GGGACGGGTG	ACTGGTGGA	ACCATTAGCT	GGGCAACAGA	
TGCCAGGATG	CCCCAGAGTG	CTCAGGGTCC	TACTGGCTGA	GTAGGAGACA	12900
CTTCGTTAAG	ACACCAGGCA	GTCCTTCCCC	TTGCTCTTCA	AATCTGAAGA	
AGTCTTTGAG	GATGGAAGAT	CATGCCCCAA	GGGCTGGCAG	CCCTTCCAAC	13000

FIGURE 1E

TCAGATATGT	AGATTCTTGG	ATCTACGATA	GCTCATTGGT	TCTAGGACAT	
ACACTCTTAT	AGCTCTGAAA	TCAAACCTCC	TATAACTGGT	GA CT CATCAT	13100
GGTTGAATTG	GCAGCTCTGT	TTGCGTCTGG	G TAGTAATGT	AAAGAAAAGT	
GCCTTTTATT	CTTGATGGCG	TCTTAGGTTT	GATGCAATAT	GGTATTTCTT	13200
CATTAGTCAC	TGTCCAGGCC	TCCTTACTCC	TGGCTCCACA	GAGGCTGTTC	
TTGTCACTCA	CTTGCAAAGA	ATAAACTCTG	AGGGCTCTCA	GAGTTTGAAC	13300
CCCAGCATAG	CCACTTACTG	GCTATGTGAC	GTTGGGCAAG	TTTCTTAACA	
TCTCTGAGCC	TGACTTTTCT	TTTGGTGTTT	TTTTTTTTTT	TTTTTTTTTTG	13400
AGACAGGGTT	TCACTCTGTC	ACCCAGGCTG	GAGTGCAGTG	GTGCAACCGT	
GGCTCAGCCT	CCACCTCCAG	GGCTCAAGCC	ATCCTCTTGC	CTTAGCCTCC	13500
TGAGTAGCTG	GGATTAGAGG	CACACACCAC	TACACCCAGC	TAATGTTTTA	
CTTTTTGTAG	AGACAGGGTC	CTACTATATT	GCCCAGGCTG	GCCTCGGACT	13600
CCTGGGCTCA	AGCGATCTTC	CGCCTCAGCC	TCCCAAAGTG	CTAGGATTAC	
GGGCATGAGC	CACCACGCCT	GGCCTGGGCC	TTAGATTTCT	TATATTTAAA	13700
GTAAGCATAA	TGACATTTCAT	TTGGTGAATT	TGTGAGAACC	AAAAACAAAG	
AAACAAACAA	AACCTACAAC	ACGTCTGACA	CAAACTATT	TATTTTCCAT	13800
TAATCTTCTT	TTTTTTTTTT	TTTTTTTTTT	TTGACACAGA	GTCTTGCTCT	
GTGCGCCAGG	CTGGAATGCA	GTGGCGCGAT	CTCGGCTCAC	TGCAACCTCT	13900
GCCTCCCAGA	TTCAAGCAAT	TCTCCTGCTT	CAGCCTCCCA	AGTAGCTGGG	
ATTACAGGCA	CGTGCCACCA	TGCCTGGCTA	ATTTTTGTAT	TTTTAGTAGA	14000
GATGGGGTTT	CACCATCTTG	GTCAGGCTGG	TCTCAAACCT	CTGGTGATCC	
ACCTGCCTCT	GCCTCCCCAA	GTGCTGGGAT	TACAGCCGTG	AGCCACTGCA	14100
CCCAGCCGGC	TTCATCTCTT	CTTGAAATCA	CTTTTATACC	ATTCTATGTG	
GTTCTCACCA	TGAGCTTGAG	TGGTGGGCTA	AAGTGCCTCT	CCCTGCTTTC	14200
AGCTTCCTGC	TGGGAACCTA	CTCTCTCAAG	TTCTTCCAG	CACCACCCCA	
TAGAGTTCCC	ATCACTCCAC	ACTGTCCAGT	GACAACCTCC	AACATGGAAG	14300
ATCTGCTAGT	TCTACAGGGT	GCTCTCTGGC	TGCCCCAGTA	ACATGTGTTT	
TTAAATTTTT	CACATGCATG	TTTGACCCCG	ACTCCCCGAA	GTCAGGTA CT	14400
GTAAGTAGCA	GTGTCAATTA	AGAAAAAGCC	CTTTAACCTC	TCTTTGCCAA	
AGGATTCTTA	TCAGCAAAAC	AGTGATGAAA	CAACAATCCC	ATAACAGCTA	14500
GCTGGCTACC	TTCTCAAGCA	CTTATTAAAT	GAGGCATAAT	GATTTTGCTT	
AATCCTCAAT	CCTGAGAGGT	GGGCGATCCC	TGTGGTGATG	AGGAAACCGA	14600
GGCTTGGGGG	TTAATGGCTT	GCCTAGATTG	ACACTGCTAG	CCAAGGAATG	
AACTGGAATT	TACACCCTGA	CCCTGACTGC	TTTTACACATT	TTCTACACAG	14700
CCTTTTCAAG	ATCCCTGCCA	ATTCTAAAAT	TAAATGATTG	TATGATTAAC	
TGTGTTTCAT	TCTTCTGCAT	CAGTTCCCAA	AACAAATTAT	ATCAAGAGAC	14800
AGCAAAAATA	TTTGTAAGA	AAGGATGTCC	AACAATCTGT	GTGGTTGTTT	
A					
TTCTGTGTTT	CTCCAATGGT	AGGGCCTCTG	TTCACCACTG	CCGTCTCTTC	14900
TTTTAGCTGT	AAGAAAAGCC	TGGAGTGCAC	GAAGTTGTGC	CTACCCCA	
[EXON 6: 14907..					
TTGAGAATGT	TAAGGGCACT	GAGGACTCAG	GTGAGGAGAA	GTGACCTGGT	15000
G					
..14980]					
GCCCATGCTC	ACCTGCCCTC	TCCCTCTTCT	TGCCCCCACC	CGTCCATCCA	
TCCCACCCAT	CCATCTATCC	CTGCGGCCCC	CCTCTGCCCC	CTCCTCTGAC	15100
T T					
CAACACCTGC	TTTGTCTGCA	GGCACCACAG	TGCTGTTGCC	CCTGGTCATT	
[EXON 7: 15122..					
TTCTTTGGTC	TTTGCCTTTT	ATCCCTCCTC	TTCATTGGTT	TAATGTATCG	15200
CTACCAACGG	TGGAAGTCCA	AGCTCTACTC	CATTGGTGAG	TGGGGGCTTT	
..15235]					
GGGAGGGAGA	GGGAGCTGGT	GGGGGTGAGG	GAGGACATGG	GTGGGTGCGA	15300
TGGACATGTG	TGGAGGGAGG	TGAGGAGTGT	CCCCTCAGTT	CATACCGCTG	
GGGACTCTGG	GCAGAAGGTG	GCCCTGGATG	GCTGGGGAGA	TGTCGAGCTG	15400
CATCAGTAGC	TCTCTCGTCC	CTGGGGCCAC	ATAGGCCCTG	AGGCATGTCA	

FIGURE 1F

CCACAAGTCC	CCACTGCCAG	CTGAGTCCAG	GGTGCCAGGG	CTGAGAGAGG	15500
AAGTGAAATT	TATGATGCTT	TCTTTCTTTT	TCCTCAGTTT	GTGGGAAATC	
C					
[EXON 8: 15538..					
GACACCTGAA	AAAGAGGTGA	GATGAAATGA	GAGAGTTACT	CCCAAATGTC	15600
..15566]					
CCTGACCATT	CCTTATAATT	GCCTAATGCT	CAGATCCCCT	GGAATCATCC	
TTCACTTTCC	GGGGGCTCGC	CTCATTCCCT	CTAAGTCCCA	ACCCCCACGT	15700
AGAATAAAGA	GGGCCGGGGC	TGGTTTTTCGC	TGCCGCACTA	ATGTGCGCCA	
CCTTCTCTCT	TTCAGGGGGA	GCTTGAAGGA	ACTACTACTA	AGCCCCCTGGC	15800
[EXON 9: 15766..					
CCCAAACCCA	AGCTTCAGTC	CCACTCCAGG	CTTCACCCCC	ACCCTGGGCT	
TCAGTCCCGT	GCCCAGTTCC	ACCTTCACCT	CCAGCTCCAC	CTATACCCCC	15900
GGTGACTGTC	CCAACCTTGC	GGCTCCCCGC	AGAGAGGTGG	CACCACCCTA	
A					
TCAGGGGGCT	GACCCCATCC	TTGCGACAGC	CCTCGCCTCC	GACCCCATCC	16000
CCAACCCCTT	TCAGAAGTGG	GAGGACAGCG	CCCACAAGCC	ACAGAGCCTA	
GACAGTGAGT	TTCTCCCGCG	GCTGGAGACG	AGGAGGCTGG	GGGAGGGCCG	16100
..16054]					
GGGGAGCGCG	GGAGGCGCTC	CCAGAGGGGA	CCACGAGAGG	CGGAGGGCGC	
GGGATGCGGG	GCGGGGCCTG	GGGTTGCCGC	CCGAGGCTCA	CCGGCCCCGC	16200
A					
TCCCCGCAGC	TGATGACCCC	GCGACGCTGT	ACGCCGTGGT	GGAGAACGTG	
[EXON 10: 16210..					
CCCCCGTTGC	GCTGGAAGGA	ATTCGTGCGG	CGCCTAGGGC	TGAGCGACCA	16300
CGAGATCGAT	CGGCTGGAGC	TGCAGAACGG	GCGCTGCCTG	CGCGAGGCGC	
AATACAGCAT	CTGGCGGACC	TGGAGGCGGC	GCACGCCGCG	GCGCGAGGCC	16400
ACGCTGGAGC	TGCTGGGACG	CGTGCTCCGC	GACATGGACC	TGCTGGGCTG	
CCTGGAGGAC	ATCGAGGAGG	CGCTTTGCGG	CCCCGCCGCC	CTCCCGCCCCG	16500
CGCCCAGTCT	TCTCAGATGA	GGCTGCGCCC	CTGCGGGCAG	CTCTAAGGAC	
..16520]					
CGTCCTGCGA	GATCGCCTTC	CAACCCCACT	TTTTTCTGGA	AAGGAGGGGT	16600
CCTGCAGGGG	CAAGCAGGAG	CTAGCAGCCG	CCTACTTGGT	GCTAACCCTT	
CGATGTACAT	AGCTTTTCTC	AGCTGCCTGC	GCGCCGCCGA	CAGTCAGCGC	16700
TGTGCGCGCG	GAGAGAGGTG	CGCCGTGGGC	TCAAGAGCCT	GAGTGGGTGG	
TTTGCGAGGA	TGAGGGACGC	TATGCCTCAT	GCCCGTTTTG	GGTGTCTCTA	16800
CCAGCAAGGC	TGCTCGGGGG	CCCCTGGTTC	GTCCCTGAGC	CTTTTTCACA	
GTGCATAAGC	AGTTTTTTTT	GTTTTTGTTT	TGTTTTTGTTT	TGTTTTTAAA	16900
TCAATCATGT	TACACTAATA	GAAACTTGGC	ACTCCTGTGC	CCTCTGCCTG	
GACAAGCACA	TAGCAAGCTG	AACTGTCCTA	AGGCAGGGGC	GAGCACGGAA	17000
CAATGGGGCC	TTCAGCTGGA	GCTGTGGACT	TTTGTACATA	CACTAAAATT	
CTGAAGTTAA	AGCTCTGCTC	TTGGAGACAG	TGGTCTGTCT	GGATGGGAGG	17100
TGGGGGCAGA	GGCCCAGATC	CTGAGGGGTG	AGATGGGAAA	AGCCCTGCAC	
TAGGGCCAGG	TAGCCCATCA	CCATCACGCC	AAGTGACAGA	GGAGTAGCAG	17200
GTTCTTGTTT	TGAACACCGT	CATCTGTTGC	CCAAGCTGGA	GTGCGCTCAC	
TGCAGCCTCC	AACCCTTGGG	CTCATGGGGT	CCTCCCGCCT	CAGCCTCCGG	17300
ACACAGGCAC	ACCACCACAC	CTGGGTAAAT	TTTAAATTTT	TTTTTTGTAA	
AGACAGGGTT	TCCCTATATT	GCCCAGGCTG	GTCTGGAAC	CCTGGGCTCA	17400
AGGGATCCTC	CCACCTCAGC	CTCCCAAAGT	GCTGGGATTA	CAGGCAGCCA	
TGCCCAGCCA	GGGCAGTCAT	TTTTATGCAC	AACTTTCTGT	GGGGCTCAGG	17500
TGCACCTATG	ATACATAAAT	TTACAGTTCT	TGATCCCCAA	ACAGAGCAGG	
AGGCAGGGTG	CCTGGGCCAG	GCTTCCTTTG	GGAAATGTGG	TCCTTGAGGT	17600
AGAGTCACAG	ATGCCGGAGG	GTGACCAGCA	CTACTGGGGA	GAGATCTCCT	
CTGGGAGAGA	TGCATGCCAA	AGGTCCTCTG	CATTTCCTCAT	ACCTCTCTGA	17700
AAAGACAGGA	GGGGGTGTTA	GGCGACATTC	AGTGGCAACG	GGTGAGGGTC	
AGGTGAAGAG	TGAGGCGGAG	AGCCCTTCCT	GCCTCAGCCC	CTGTTCTCTG	17800

FIGURE 1G

TTTGCCCTCT	TTCTATACTA	CACCCACCA	CCATACAGAC	ATCCCCGTCT	
GCCCCCTCCC	AGGCCAGCTT	CCCTCCAGCA	CTTACGATGC	GGACAGAGGG	17900
GTGTCCAGCT	GAATGATGTG	GGGCCCCCGC	ATCCTCTGCA	GCTGGGCCCCG	
AGTCAGCTTC	CGTGGCCTGC	TGTCCCGGGG	CTCCTCGGCC	CCCTCAATCC	18000
TTTGGCTGGC	CAGCTCCTCC	CGGATCTCTC	TGAGCATGTC	CTCAGCCCCGC	
ATTGGGCGCA	GGGATGTGTG	GCCAGCTTTC	AGGAACAGAG	GCCCCCTCTTC	18100
TTCTCCTCC	CCTGAGGACT	CCCAGGGGCT	TTCCCCGGCA	GAGTCAGCAT	
GGGTGGGGGA	GGAGGGAAGC	TGGCCCCGAA	GCCGGGCCCT	GTGGAGTGTT	18200
TCCACCACCA	CATTCCCTCG	CTCGGAGGCC	CCATCTTCTT	CCTCAGACCA	
GGTTGGTGGG	TCTTCCTGGG	GAAGACTGCC	TCCTTTTAGG	ATTCTTTCCG	18300
GCAGTTCGGG	GGCGCTTCGG	CGTTGAGGAG	CTTGGGGGTC	GGGAGGGTGG	
GGACGCAGAG	GGATGTCCCG	GAGTTCAGG	GTGGAGAAGG	TGAGGCGAGG	18400
GTCCCGCCGA	AGGGCTCTTT	GGCGTAGACG	GCTCAGTGGG	GAGCGGGACC	
CCGTGGGGGT	GCCTGGGATC	AAAGTGCCGT	AGCCAGAGTC	TGAGGTATCA	18500
TCTGGCACAA	GGGGAGCATC	TTCATCTGTG	TCTTCTGTCA	CCACCAGGTG	
GGGGATAATG	GTCGAGAACT	CAGGAGTCCT	ACAGTTAATG	GCAAAGAGTC	18600
AGATGCGTAG	GGGTCAAGTT	CAAGTCCAGG	GAGTTTCCCT	TGATCACTAC	
ATCCAGAAAT	GGCCCCCTCT	CCAAACTTAT	TTTGGTATCA	TCTTTCCATC	18700
GCACTGTGAT	TGTTTTTCTC	ATCTGGCTGG	CTAGATTTTA	AGCTCCTAAG	
AGAGTACGGG	CTGCCTCTAT	ACTGTTTTAT	CCATAGCATC	TGGTCCAGGA	18800
TCTTGTATCG	AGTGGGTAGT	CAGGTTTTTG	CTGAGTGGTT	CCTGAACTTA	
CCTGATATTA	TCCTCAATGA	TCGATTCTTC	TTTTCTCCTT	AAGCTGCTGC	18900
CAAGCAGTGG	TGCTATCCTA	GACGAACCTC	ACACTCCCCG	GGGATTTGGC	
AGCTCTAATA	TTCTGCAGAT	CCACACCTAC	CTTCACTCTC	GAGCTTGCTC	19000
CTCTCACAGT	GCTCCTGTGT	GACTCTAGGC	AGGCTAACTC	TGTAGGCTGT	
CTGTGCCCTA	TCCCCACCT	CCAACCCAAC	ACGGCTGGTA	CCAACCTTCC	19100
GACCCAACAC	AGCTGGTACC	GAGCTTCCCT	ACCCTGCCCT	ACGCCTGCGT	
TCCTCTATCT	ATTCCCAATT	CCACCAAAAA	TGTGCAGTAA	TGCCATTTCT	19200
CAGCCTTATG	GCTCCCTCCT	CCTGCTCGGG	GAGACCTTGT	AGTCCGTGTG	
AGCCTTACCT	CCCCTCTGCG	CTGCTCTGAG	AGCCCTCCAG	GGAAGGCGTG	19300
GAGGGCCTGG	TGCTGGGGGA	CTCCCTGTCC	TGGTCCCGAT	AGAGGGTCAG	
GAGCTCCCTC	TTCTGTTGAA	CATACTCCTC	TGCCTTCAGC	TTCTGTAGGG	19400
CGGCCTGGGA	CAGGACACTT	TCGTTATTAA	GAGCTCTCAT	TTATTGAGCA	
CTTGCTGTTT	GCCAGGCACC	CTGCTAAGTG	CGTTACATAT	ATTACCTTAT	19500
TTTATTTTAT	TATTATTATT	ATTTTTTGAG	ACTGAGTCTT	GCTCTGTCAC	
CCAGACTAGA	GTGCAGTGCC	ACAATCTTGG	CTCACTGCAA	CCTCCACCTC	19600
CTGGGTTCAA	GCGATTCTCC	TGCCTCAGCC	TCCTTAGTAG	CTGGGATTAC	
AGGCGCCCGC	CAACGTGCCC	GGCTAATTTT	TGTATTTTTA	GTAGAGATGG	19700
GGTTTCACCA	TCTTGGCCAG	GCTGGTCTCA	AACTCCTGAC	CTTGTGATCC	
ACCCCCCTTG	GCCTCCCAAA	GTGCTGGAAT	TAGACGTGTA	AGCCACCGTG	19800
CCCGGCCTAC	ATTACCTTAT	TTAATCTTTA	CAAAAACCCC	ATGAACCAGA	
TATTTTACC	CCACCTTACT	ACTGAGACAT	GGAGACTCTA	AGGTAAAGTA	19900
ACTGTCTGAG	GGGGTACTTC	TTACCATAAG	AAAGTGGGGT	GGTGCCGGGA	
TTTGGTGGCA	CCAAACTCTG	GAGCTAGTGT	TGGGGGTGAG	TGGGGTGAAC	20000
AGAATGGCCC	TTTTCTTACC	TGTACAGGTC	TTCTTGCTTC	TCATGTCCCA	
TTGGCAGACC	TGTTATCAGG	TCTTCCCCCT	CCTTCAGGAA	GCCCTCCCTG	20100
GTTGGTGGTG	ATGGTAGAAT	AAGTGTTCTG	AATTGGTACT	GGTTGCTCCT	
TCAAGAGCAT	CCCTCTCCTA	CCACCTGGGC	CTCTGCCCTG	AAGCTGGGAG	20200
GAGCAGGAGG	GCAGAACGTG	GGCAGAGGTG	GGCTTTGTCC	CAGGCTGAGG	
ACTCTGCTGT	CCTTCAGAGG	GAGGAAAGTT	CCTAGAAGGC	TGAGGAGAGG	20300
ACGCATTATA	TTATCTGCCT	TCTCCCTCCC	TCAGCGATTT	CATACAGGTA	
CCATCAAAAG	GAAATAGCGC	CACCTGAGAA	AAAATTTTCA	AAGCACTTTT	20400
GCACATGTGG	TCATTTGATA	CACATCATTG	CCCTGTGGTG	TGGAGAACAT	
GAATGTTAGC	CCATTTTACA	GACAAGAAAC	CTAGACCTAG	AGAGGTGAAG	20500
TGACTTGCTC	AAGGTGCCA				20519

FIGURE 1H

# POLYMORPHISMS IN THE CODING SEQUENCE OF TNFRSF1A

ATGGGCCTCT	CCACCGTGCC	TGACCTGCTG	CTGCCCCAGG	TGCTCCTGGA	
GCTGTTGGTG	GGAATATAACC	CCTCAGGGGT	TATTGGACTG	GTCCCTCACC	100
TAGGGGACAG	GGAGAAGAGA	GATAGTGTGT	GTCCCCAAGG	AAAATATATC	
CACCCTCAAA	ATAATTTCGAT	TTGCTGTACC	AAGTGCCACA	AAGGAACCTA	200
CTTGTACAAT	GACTGTCCAG	GCCCGGGGCA	GGATACGGAC	TGCAGGGAGT	
		T			
GTGAGAGCGG	CTCCTTCACC	GCTTCAGAAA	ACCACCTCAG	ACACTGCCTC	300
AGCTGCTCCA	AATGCCGAAA	GGAAATGGGT	CAGGTGGAGA	TCTCTTCTTG	
CACAGTGGAC	CGGGACACCG	TGTGTGGCTG	CAGGAAGAAC	CAGTACCGGC	400
	A				
ATTATTGGAG	TGAAAACCTT	TTCCAGTGCT	TCAATTGCAG	CCTCTGCCTC	
	C				
AATGGGACCG	TGCACCTCTC	CTGCCAGGAG	AAACAGAACA	CCGTGTGCAC	500
CTGCCATGCA	GGTTTCTTTC	TAAGAGAAAA	CGAGTGTGTC	TCCTGTAGTA	
ACTGTAAGAA	AAGCCTGGAG	TGCACGAAGT	TGTGCCTACC	CCAGATTGAG	600
AATGTTAAGG	GCACTGAGGA	CTCAGGCACC	ACAGTGCTGT	TGCCCCCTGGT	
CATTTTCTTT	GGTCTTTGCC	TTTTATCCCT	CCTCTTCATT	GGTTTAATGT	700
ATCGCTACCA	ACGGTGGAAG	TCCAAGCTCT	ACTCCATTGT	TTGTGGGAAA	
TCGACACCTG	AAAAAGAGGG	GGAGCTTGAA	GGAACACTA	CTAAGCCCCT	800
GGCCCCAAAC	CCAAGCTTCA	GTCCCCTCC	AGGCTTCACC	CCCACCCTGG	
GCTTCAGTCC	CGTGCCCAGT	TCCACCTTCA	CCTCCAGCTC	CACCTATACC	900
CCCGGTGACT	GTCCCAACTT	TGCGGCTCCC	CGCAGAGAGG	TGGCACCACC	
		A			
CTATCAGGGG	GCTGACCCCA	TCCTTGCGAC	AGCCCTCGCC	TCCGACCCCA	1000
TCCCCAACCC	CCTTCAGAAG	TGGGAGGACA	GCGCCACAA	GCCACAGAGC	
CTAGACACTG	ATGACCCCGC	GACGCTGTAC	GCCGTGGTGG	AGAACGTGCC	1100
CCCGTTGCGC	TGGAAGGAAT	TCGTGCGGCG	CCTAGGGCTG	AGCGACCACG	
AGATCGATCG	GCTGGAGCTG	CAGAACGGGC	GCTGCCTGCG	CGAGGCGCAA	1200
TACAGCATGC	TGGCGACCTG	GAGGCGGCGC	ACGCCGCGGC	GCGAGGCCAC	
GCTGGAGCTG	CTGGGACGCG	TGCTCCGCGA	CATGGACCTG	CTGGGCTGCC	1300
TGGAGGACAT	CGAGGAGGCG	CTTTGCGGCC	CCGCCGCCCT	CCCGCCCGCG	
CCCAGTCTTC	TCAGATGA				1368

0945505:083404

FIGURE 2

# ISOFORMS OF THE TNFRSF1A PROTEIN

MGLSTVPDLL	LPQVLLELLV	GIYPSGVIGL	VPHLGDREKR	DSVCPQGKYI	
HPQNNsicCT	KCHKGTyLYN	DCPGPGQDtd	CRECESGSFT	ASENHLRHCL	100
		L			
SCSKCRKEMG	QVEISSCTVD	RDtVCGCRKN	QYRHYWSENl	FQCFNCSLCL	
		Q	H		
NGTVHLSCQE	KQNTVCTCHA	GFFLRENECV	SCSNCKKSLE	CTKLCLPQIE	200
NVKGTEdSGT	TVLLPLViff	GLCLLSLLFI	GLMYRYQRWK	SKLYSIVCGK	
STPEKEGELE	GTTTKPLAPN	PSFSPTPGFT	PTLGFSPVPS	STFTSSSTYT	300
PGDCPNFAAP	RREVAPPYQG	ADpILATALA	SDPIPNPLQK	WEDSAHKPQS	
	K				
LDTDDPATLY	AVVENVPPLR	WKEFVRRlGL	SDHEIDRLEL	QNGRCLREAQ	400
YSMLATWRRR	TPRREATLEL	LGRVLRDMDL	LGcLEDIEEA	LCGPAALPPA	
PSLLR					455

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TOTED 5054660

FIGURE 3